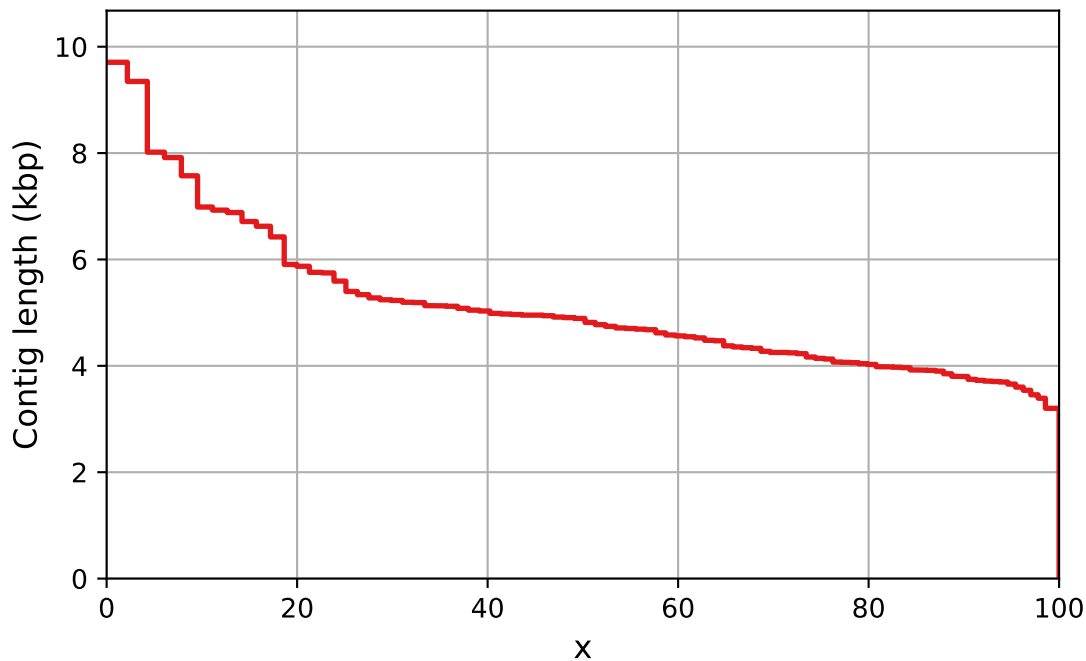


Report

	genome.nextpolish
# contigs (>= 0 bp)	92
# contigs (>= 1000 bp)	92
# contigs (>= 5000 bp)	29
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	445780
Total length (>= 1000 bp)	445780
Total length (>= 5000 bp)	179398
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	92
Largest contig	9707
Total length	445780
GC (%)	54.67
N50	4891
N90	3798
auN	5153.8
L50	38
L90	80
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 0 part

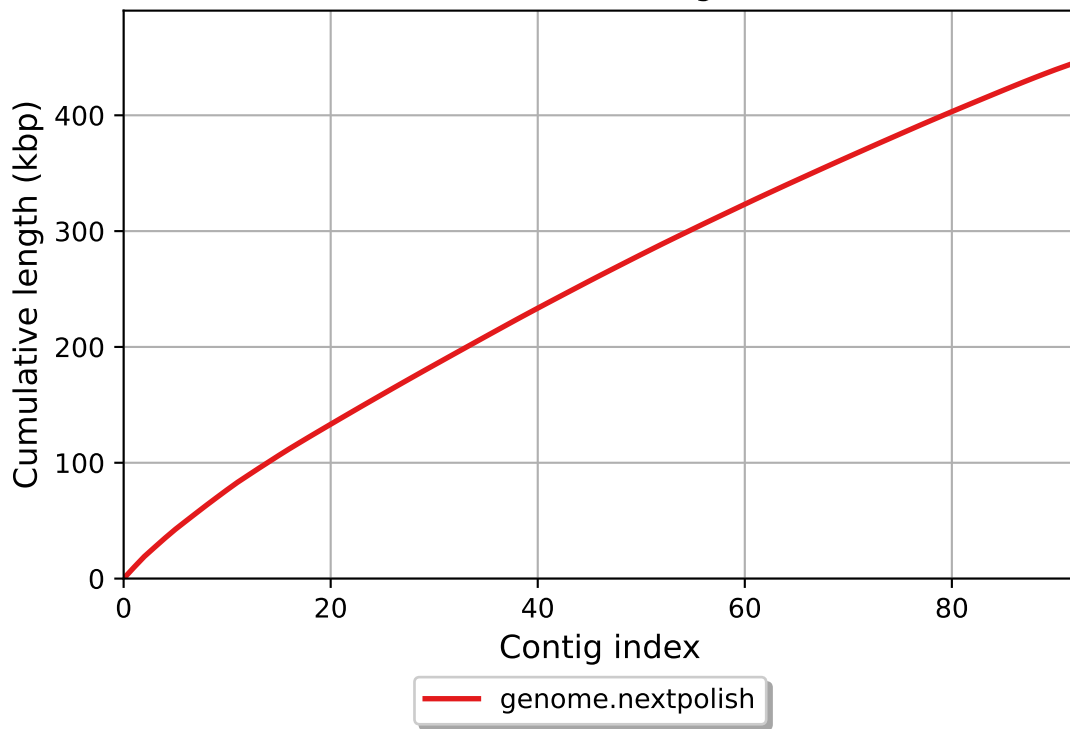
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

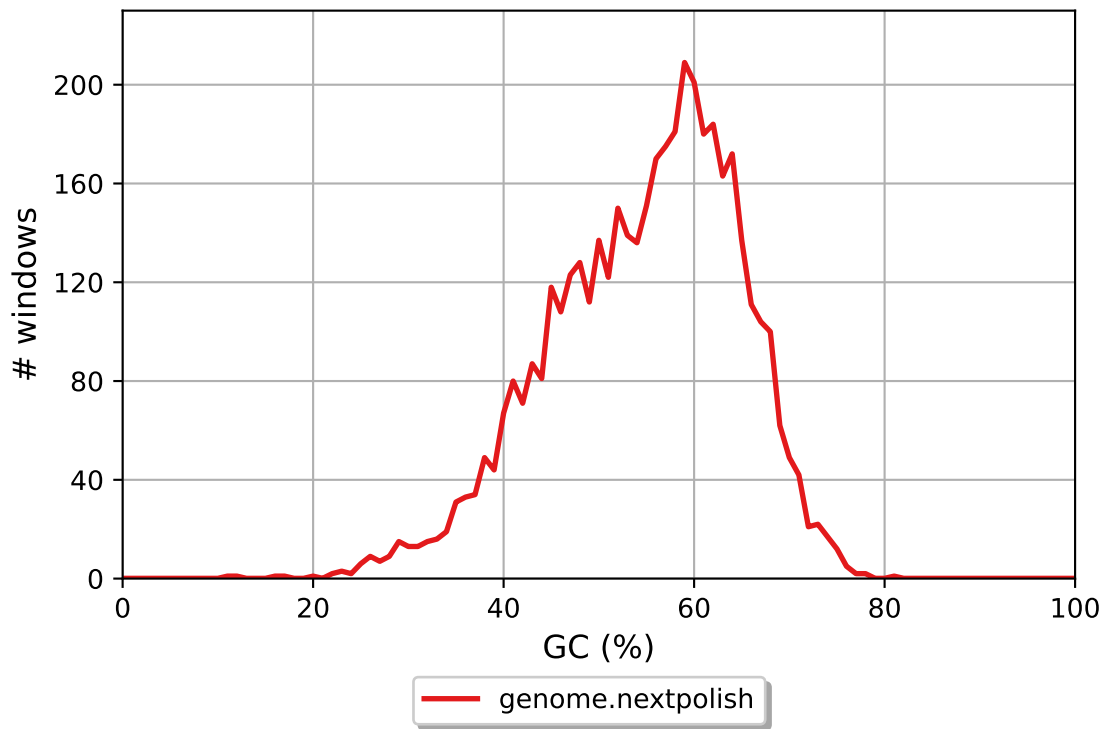


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

